SEQUENCE LISTING

(1) GENERAL INFORMATION:

   (i) APPLICANT: HAWLEY-NELSON, PAMELA
       LAN, JIANQING
       SHIH, POJEN
       JESSE, JOEL A.
       SCHIFFERLI, KEVIN P.
       GEBEYEHU, GULILAT

   (ii) TITLE OF INVENTION: PEPTIDE-ENHANCED TRANSFECTIONS

   (iii) NUMBER OF SEQUENCES: 120

   (iv) CORRESPONDENCE ADDRESS:
      (A) ADDRESSEE: GREENLEE, WINNER & SULLIVAN
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      (D) STATE: CO
      (E) COUNTRY: US
      (F) ZIP: 80303

   (v) COMPUTER READABLE FORM:
      (A) MEDIUM TYPE: Floppy disk
      (B) COMPUTER: IBM PC compatible
      (C) OPERATING SYSTEM: PC-DOS/MS-DOS
      (D) SOFTWARE: PatentIn Release #1.0, Version #1.30

   (vi) CURRENT APPLICATION DATA:
      (A) APPLICATION NUMBER: Not assigned
      (B) FILING DATE: 23-JUL-2001
      (C) CLASSIFICATION:

   (vii) PRIOR APPLICATION DATA
      (A) APPLICATION NUMBER: US 09/039,780
      (B) FILING DATE: 16-MAR-1998
      (C) CLASSIFICATION:

   (viii) ATTORNEY/AGENT INFORMATION:
      (A) NAME: SULLIVAN, SALLY A.
      (B) REGISTRATION NUMBER: 32,064
      (C) REFERENCE/DOCKET NUMBER: 32-95D

   (ix) TELECOMMUNICATION INFORMATION:
      (A) TELEPHONE: (303)499-8080
      (B) TELEFAX: (303)499-8089

(2) INFORMATION FOR SEQ ID NO:1:

    (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 10 amino acids
       (B) TYPE: amino acid
       (C) STRANDEDNESS: not relevant
       (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

    His Ala Ile Arg Gly Asp Thr Phe Ala Thr
    1   5   10

(2) INFORMATION FOR SEQ ID NO:2:

    (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 7 amino acids
       (B) TYPE: amino acid
       (C) STRANDEDNESS: not relevant
       (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

    Pro Lys Lys Lys Arg Lys Val
    1   5

(2) INFORMATION FOR SEQ ID NO:3:

    (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 8 amino acids
       (B) TYPE: amino acid
       (C) STRANDEDNESS: not relevant
       (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

Pro Lys Lys Lys Lys Arg Lys Val
1   5

(2) INFORMATION FOR SEQ ID NO:4:

    (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 20 amino acids
       (B) TYPE: amino acid
       (C) STRANDEDNESS: not relevant
       (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (ix) FEATURE:
       (A) NAME/KEY: Modified-site
       (B) LOCATION: 2..20
       (D) OTHER INFORMATION: /product= "OTHER"
/note= "ANY OR ALL OF THE AMINO ACIDS 2-20 CAN BE PRESENT OR
ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Lys Lys Lys Lys
            20

(2) INFORMATION FOR SEQ ID NO:5:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 2..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "ANY OR ALL OF THE AMINO ACIDS 2-20 CAN BE PRESENT OR
ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

    Arg Arg Arg Arg Arg Arg Arg Arg Arg Arg Arg Arg Arg Arg Arg Arg
    1   5   10   15

    Arg Arg Arg Arg
            20

(2) INFORMATION FOR SEQ ID NO:6:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 40 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 3..40
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "ANY Lsy-Arg PAIR FROM AMINO ACID 3-40 CAN BE PRESENT OR
ABSENT"

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg
1   5   10   15

Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg Lys Arg
    20   25   30

Lys Arg Lys Arg Lys Arg Lys Arg
    35   40

(2) INFORMATION FOR SEQ ID NO:7:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 2..20
        (D) OTHER INFORMATION: /product= "OTHER"
    /note= "ANY OR ALL OF THE AMINO ACIDS 2-20 CAN BE PRESENT OR
    ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

    Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly
    1   5   10   15

    Gly Gly Gly Gly
            20

(2) INFORMATION FOR SEQ ID NO:8:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 7 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

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(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Gly Arg Gly Asp Ser Pro Cys
1   5

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(2) INFORMATION FOR SEQ ID NO:9:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 23 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "ANY OR ALL OF THE AMINO ACIDS 1-20 CAN BE PRESENT OR
ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Lys Lys Lys Lys Arg Gly Asp
     20

(2) INFORMATION FOR SEQ ID NO:10:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 19 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Arg Gly Asp

(2) INFORMATION FOR SEQ ID NO:11:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 7 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

    Lys Phe Thr Ile Val Phe Cys
    1   5

(2) INFORMATION FOR SEQ ID NO:12:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 12 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

Cys Gly Trp Gly Pro Lys Lys Lys Arg Lys Val Gly
1   5   10

(2) INFORMATION FOR SEQ ID NO:13:

   (i) SEQUENCE CHARACTERISTICS:
      (A) LENGTH: 26 amino acids
      (B) TYPE: amino acid
      (C) STRANDEDNESS: not relevant
      (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (iii) HYPOTHETICAL: NO

   (iv) ANTI-SENSE: NO

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

Gly Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly Gly Gly Tyr
1   5   10   15

Gly Pro Lys Lys Lys Arg Lys Val Gly Gly
      20   25

(2) INFORMATION FOR SEQ ID NO:14:

   (i) SEQUENCE CHARACTERISTICS:
      (A) LENGTH: 6 amino acids
      (B) TYPE: amino acid
      (C) STRANDEDNESS: not relevant
      (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (iii) HYPOTHETICAL: NO

   (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Arg Gly Asp Ser Pro Cys
1   5

(2) INFORMATION FOR SEQ ID NO:15:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 4 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (iii) HYPOTHETICAL: NO

   (iv) ANTI-SENSE: NO

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

Arg Glu Asp Val
1

(2) INFORMATION FOR SEQ ID NO:16:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 4 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (iii) HYPOTHETICAL: NO

   (iv) ANTI-SENSE: NO

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Arg Gly Asp Val
1

(2) INFORMATION FOR SEQ ID NO:17:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 25 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

Asp Glu Leu Pro Gln Leu Val Thr Leu Pro His Pro Asn Leu His Gly
1   5   10   15

Pro Glu Ile Leu Asp Val Pro Ser Thr
      20   25

(2) INFORMATION FOR SEQ ID NO:18:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 4 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Arg Gly Asp Asn
1

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(2) INFORMATION FOR SEQ ID NO:19:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 4 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

    Arg Gly Asp Met
    1

(2) INFORMATION FOR SEQ ID NO:20:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 4 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

    Arg Gly Asp Thr
    1

(2) INFORMATION FOR SEQ ID NO:21:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 4 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (iii) HYPOTHETICAL: NO

   (iv) ANTI-SENSE: NO

   (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 4
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 4 CAN BE ANY AMINO ACID"

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

   Arg Gly Asp Xaa
   1

(2) INFORMATION FOR SEQ ID NO:22:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 4 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (iii) HYPOTHETICAL: NO

   (iv) ANTI-SENSE: NO

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

   Arg Gly Asp Ser
   1

(2) INFORMATION FOR SEQ ID NO:23:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 4 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (iii) HYPOTHETICAL: NO

   (iv) ANTI-SENSE: NO

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

   Arg Gly Asp Ala
   1

(2) INFORMATION FOR SEQ ID NO:24:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 33 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "C AT POSITION 1 CAN BE ABSENT"

   (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 3
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 3 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 14..33
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "ANY OR ALL OF THE GLY AT POSITIONS 14 TO 33 CAN BE
ABSENT"

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

Cys-Gly-Xaa-Gly-Pro-Lys-Lys-Lys-Arg-Lys-Val-Gly-Gly-Gly-Gly-Gly
1   5   10   15

Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly Gly
      20   25   30

Gly

(2) INFORMATION FOR SEQ ID NO:25:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 48 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS AND ANY OR
ALL OF THESE AMINO ACIDS CAN BE ABSENT "

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 28
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 28 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 29..48
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 29-48 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys Val Xaa Xaa Xaa Xaa Xaa
     20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   35   40   45

(2) INFORMATION FOR SEQ ID NO:26:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 90 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR ANY OF
THESE AMINO ACIDS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITON 21 CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 22 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 23..42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 23-42 CAN BE ANY AMINO ACIDS OR ANY OF
THESE AMINO ACIDS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 50 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 51..70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 51 TO 79 CAN BE ANY AMINO ACIDS OR ANY
OF THESE AMINO ACIDS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 71..90
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 71-90 CAN BE ANY CATIONIC AMINO ACIDS
OR ANY OF THESE AMINO ACIDS CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
    35   40   45

Val Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
           85   90

(2) INFORMATION FOR SEQ ID NO:27:

   (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 70 amino acids
       (B) TYPE: amino acid
       (C) STRANDEDNESS: not relevant
       (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (ix) FEATURE:
       (A) NAME/KEY: Modified-site
       (B) LOCATION: 1
       (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

   (ix) FEATURE:
       (A) NAME/KEY: Modified-site
       (B) LOCATION: 2..21
       (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR ANY OF
THESE AMNION ACIDS CAN BE ABSENT"

   (ix) FEATURE:
       (A) NAME/KEY: Modified-site
       (B) LOCATION: 22
       (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 22 CAN BE TYR OR TRP OR CAN BE ABSENT"

   (ix) FEATURE:
       (A) NAME/KEY: Modified-site
       (B) LOCATION: 23..42
       (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 23 TO 42 CAN BE ANY AMNIO ACIDS OR ANY
OF THESE AMINO ACIDS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50 TO 69 CAN BE ANY AMINO ACIDS OR ANY
OF THESE AMINO ACIDS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 70 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
    35   40   45

Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

Xaa Xaa Xaa Xaa Xaa Cys
65   70

(2) INFORMATION FOR SEQ ID NO:28:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 90 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 2..21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR ANY OF
THESE AMINO ACIDS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 22 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 23..42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 23-42 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 50 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 51..70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 51 TO 70 CAN BE ANY AMINO ACIDS AND CAN
BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 71..90
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 71-90 CAN BE ANY CATIONIC AMINO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     20   25   30

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
      35   40   45

Val Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      85   90

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(2) INFORMATION FOR SEQ ID NO:29:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 90 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21..40
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21 TO 40 CAN BE ANY AMINO ACIDS OR CAN
BE ABSENT"

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 42 CAN BE TYR OT TRP OR CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 43..62
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 43 TO 62 CAN BE ANY AMINO ACIDS OR CAN
BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 70..89
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 70 TO 89 CAN BE ANY AMINO ACIDS OR CAN
BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 90
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 90 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys
  50   55   60

Lys Lys Arg Lys Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
    85   90

(2) INFORMATION FOR SEQ ID NO:30:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 90 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACID OR
CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21..40
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21 TO 40 CAN BE ANY AMINO ACIDS OR CAN
BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 42 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 43..62
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 43 TO 62 CAN BE ANY AMINO ACIDS OR CAN
BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 70 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 71..90
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 71 TO 90 CAN BE ANY AMINO ACIDS OR CAN
BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys
      50   55   60

Lys Lys Arg Lys Val Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      85   90

(2) INFORMATION FOR SEQ ID NO:31:

   (i) SEQUENCE CHARACTERISTICS:
      (A) LENGTH: 5 amino acids
      (B) TYPE: amino acid
      (C) STRANDEDNESS: not relevant
      (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:

Arg Gly Asp Ser Pro
1   5

(2) INFORMATION FOR SEQ ID NO:32:

   (i) SEQUENCE CHARACTERISTICS:
      (A) LENGTH: 69 amino acids
      (B) TYPE: amino acid
      (C) STRANDEDNESS: not relevant
      (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21..40
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 42 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 48
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 48 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 49 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Gly Arg Gly Asp Ser Pro Gly
    35   40   45

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

Xaa Xaa Xaa Xaa Xaa
65

(2) INFORMATION FOR SEQ ID NO:33:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 69 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 22
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 22 CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 28
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 28 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 29
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 29 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30..49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 30-49 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50 -69 CAN BE ANY CATIONIC AMINO ACIDS"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Gly Arg Gly Asp Ser Pro Gly Cys Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa Xaa
65

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(2) INFORMATION FOR SEQ ID NO:34:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 69 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21..40
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 41
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 42
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 42 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 48
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 48 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 49..68
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 49-68 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 69 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Gly Arg Gly Asp Ser Pro Gly
   35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa Cys
65

(2) INFORMATION FOR SEQ ID NO:35:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 69 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 2..21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACID OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 22 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 28
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 28 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 29
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 29 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30..49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 30-49 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY CATIONIC AMINO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Gly Arg Gly Asp Ser Pro Gly Cys Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

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Xaa Xaa Xaa Xaa Xaa
65

(2) INFORMATION FOR SEQ ID NO:36:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 5 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:

Arg Gly Asp Met Phe
1   5

(2) INFORMATION FOR SEQ ID NO:37:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 49 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 1..20
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

   (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 21
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 22 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 28
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 28 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 29
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 29 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30..49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 30-49 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Gly Arg Gly Asp Met Phe Gly Cys Xaa Xaa Xaa
        20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        35   40   45

Xaa

(2) INFORMATION FOR SEQ ID NO:38:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 47 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 2..21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 27..46
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 27-46 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 47
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 47 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Arg Gly Asp Met Phe Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
    35   40   45

(2) INFORMATION FOR SEQ ID NO:39:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 69 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

123

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21..40
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 42 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 48
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 48 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 49..68
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 49-68 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 69 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Gly Arg Gly Asp Met Phe Gly
      35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      50   55   60

Xaa Xaa Xaa Xaa Cys
65

(2) INFORMATION FOR SEQ ID NO:40:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 69 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 2..21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 22
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 22 CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 28
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 28 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 29
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 29 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30..49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 30-49 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY CATIONIC AMINO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Gly Arg Gly Asp Met Phe Gly Cys Xaa Xaa Xaa
      20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

Xaa Xaa Xaa Xaa Xaa
65

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(2) INFORMATION FOR SEQ ID NO:41:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 69 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21..40
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 41
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 42
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 42 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 48
        (D) OTHER INFORMATION: /product= "OTHRE"
/note= "GLY AT POSITION 48 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 49
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 49 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Gly Arg Gly Asp Met Phe Gly
   35   40   45

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 50   55   60

Xaa Xaa Xaa Xaa Xaa
65

(2) INFORMATION FOR SEQ ID NO:42:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 69 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 22 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 28
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "GLY AT POSITION 28 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 29
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 29 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30..49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 30-49 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY CATIONIC AMINO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Gly Arg Gly Asp Met Phe Gly Cys Xaa Xaa Xaa
      20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     50   55   60

Xaa Xaa Xaa Xaa Xaa
65

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(2) INFORMATION FOR SEQ ID NO:43:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 67 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 2..21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 47..66
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 47-66 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 67
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 67 CAN BE ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Asp Glu Leu Pro Gln Leu Val Thr Leu Pro His
      20   25   30

Pro Asn Leu His Gly Pro Glu Ile Leu Asp Val Pro Ser Thr Xaa Xaa
      35   40   45

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
50   55   60

Xaa Xaa Cys
65

(2) INFORMATION FOR SEQ ID NO:44:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 87 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not-relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACID OR
CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21..40
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 41
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 67..86
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 67-86 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 87
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 87 CAN BE ABSENT"

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Asp Glu Leu Pro Gln Leu Val
    35   40   45

Thr Leu Pro His Pro Asn Leu His Gly Pro Glu Ile Leu Asp Val Pro
    50   55   60

Ser Thr Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Cys
        85

(2) INFORMATION FOR SEQ ID NO:45:

  (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 87 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 1..20
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

  (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 21..40
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSTION 41 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 67
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 67 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 68..87
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 68-87 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Asp Glu Leu Pro Gln Leu Val
    35   40   45

Thr Leu Pro His Pro Asn Leu His Gly Pro Glu Ile Leu Asp Val Pro
   50   55   60

Ser Thr Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      85

(2) INFORMATION FOR SEQ ID NO:46:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 87 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 2..21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACID OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 47
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 47 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 48..67
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 48-67 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 68..87
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 68-87 CAN BE ANY CATIONIC AMINO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Asp Glu Leu Pro Gln Leu Val Thr Leu Pro His
     20   25   30

Pro Asn Leu His Gly Pro Glu Ile Leu Asp Val Pro Ser Thr Cys Xaa
    35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa
         85

(2) INFORMATION FOR SEQ ID NO:47:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 87 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 47
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 47 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 48..67
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 48-67 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 68..87
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 68-87 CAN BE ANY CATIONIC AMNIO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Asp Glu Leu Pro Gln Leu Val Thr Leu Pro His
    20   25   30

Pro Asn Leu His Gly Pro Glu Ile Leu Asp Val Pro Ser Thr Cys Xaa
    35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    85

(2) INFORMATION FOR SEQ ID NO:48:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 8 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:48:

Glu Ile Leu Asp Val Pro Ser Thr
1   5

(2) INFORMATION FOR SEQ ID NO:49:

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(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 50 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30..49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 30-49 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 50 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Glu Ile Leu Asp Val Pro Ser Thr Xaa Xaa Xaa
        20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        35   40   45

Xaa Cys
    50

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(2) INFORMATION FOR SEQ ID NO:50:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 70 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

(ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21..40
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 41
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 50
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 50 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 51..70
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 51-70 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
　　　　20　　　　　　　25　　　　　　　30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Glu Ile Leu Asp Val Pro Ser
　　　35　　　　　　40　　　　　　　45

Thr Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
　　50　　　　　　　55　　　　　　　60

Xaa Xaa Xaa Xaa Xaa Xaa
65　　　　　　70

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(2) INFORMATION FOR SEQ ID NO:51:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 70 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21..40
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

  (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 70 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:51:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Glu Ile Leu Asp Val Pro Ser
   35   40   45

Thr Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 50   55   60

Xaa Xaa Xaa Xaa Xaa Cys
65   70

(2) INFORMATION FOR SEQ ID NO:52:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 70 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

 (ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 2..21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 30 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 31..50
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 31-50 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 51..70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 51-70 AN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:52:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Glu Ile Leu Asp Val Pro Ser Thr Cys Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa
65   70

(2) INFORMATION FOR SEQ ID NO:53:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 70 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 50 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 31..50
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 31-50 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 51..70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 51-70 CAN BE ANY CATIONIC AMINO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:53:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Glu Ile Leu Asp Val Pro Ser Thr Cys Xaa Xaa
     20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa
65   70

(2) INFORMATION FOR SEQ ID NO:54:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 6 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:54:

Lys Phe Thr Ile Val Phe
1   5

(2) INFORMATION FOR SEQ ID NO:55:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 48 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 1
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

   (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 2..21
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

   (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 28..47

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(D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 28-47 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 48
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 48 CAN BE ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:

    Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    1   5   10   15

    Xaa Xaa Xaa Xaa Xaa Lys Phe Thr Ile Val Phe Xaa Xaa Xaa Xaa Xaa
            20   25   30

    Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
        35   40   45

(2) INFORMATION FOR SEQ ID NO:56:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 68 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACIDS OR
CAN BE AB..."

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21..40
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 48
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 48 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 49..68
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 49-68 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:56:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Lys Phe Thr Ile Val Phe Cys
   35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa
65

(2) INFORMATION FOR SEQ ID NO:57:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 68 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY CATIONIC AMINO ACID OR
CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21..40
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 21-40 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 41 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 48..67
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 48-67 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 68
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 68 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:57:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Lys Phe Thr Ile Val Phe Xaa
    35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   50   55   60

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Xaa Xaa Xaa Cys
65

(2) INFORMATION FOR SEQ ID NO:58:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 68 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 1 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 2..21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 2-21 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 28
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 28 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 29..48
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 29-48 CAN BE ANY AMNINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 49..68
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 49-68 CAN BE ANY CATIONIC AMINO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:58:

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Lys Phe Thr Ile Val Phe Cys Xaa Xaa Xaa Xaa
        20   . 25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        50   55   60

Xaa Xaa Xaa Xaa
65

(2) INFORMATION FOR SEQ ID NO:59:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 68 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 28
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 28 CAN BE ABSENT" ·

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 29..48
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 29-48 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 49..68
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 49-68 CAN BE ANY CATIONIC AMINO ACIDS
OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:59:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Lys Phe Thr Ile Val Phe Cys Xaa Xaa Xaa Xaa
     20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa
65

(2) INFORMATION FOR SEQ ID NO:60:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 95 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 22 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 23..42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 23-42 ANC BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 75
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 75 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 76..95
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 76-95 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:60:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      20   25   30

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
　　35　　　　　　40　　　　　45

Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
　　50　　　　　55　　　　　60

Xaa Xaa Xaa Xaa Xaa Arg Gly Asp Met Phe Cys Xaa Xaa Xaa Xaa Xaa
　65　　　　　70 -　　　75　　　　80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
　　　85　　　　　90　　　　95

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(2) INFORMATION FOR SEQ ID NO:61:

　(i) SEQUENCE CHARACTERISTICS:
　　(A) LENGTH: 95 amino acids
　　(B) TYPE: amino acid
　　(C) STRANDEDNESS: not relevant
　　(D) TOPOLOGY: linear

　(ii) MOLECULE TYPE: peptide

　(ix) FEATURE:
　　(A) NAME/KEY: Modified-site
　　(B) LOCATION: 1..20
　　(D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

　(ix) FEATURE:
　　(A) NAME/KEY: Modified-site
　　(B) LOCATION: 21
　　(D) OTHER INFORMATION: /product= "OTHER"
/note="CYS AT POSITION 21 CAN BE ABSENT"

　(ix) FEATURE:
　　(A) NAME/KEY: Modified-site
　　(B) LOCATION: 22..41
　　(D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 22-41 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

　(ix) FEATURE:
　　(A) NAME/KEY: Modified-site
　　(B) LOCATION: 42
　　(D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 42 CAN BE TYR OR TRP OR CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 75
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 75 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 76..95
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 76-95 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:61:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
   35   40   45

Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 50   55   60

Xaa Xaa Xaa Xaa Xaa Arg Gly Asp Met Phe Cys Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    85   90   95

(2) INFORMATION FOR SEQ ID NO:62:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 96 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

· (ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 22 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 23..42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 23-42 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OF CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 76
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 76 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 77..96
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 77-96 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:62:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   .   15

Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
        35   40   45

Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50 ·   55   60

Xaa Xaa Xaa Xaa Xaa Gly Arg Gly Asp Ser Pro Cys Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        85   90   95

(2) INFORMATION FOR SEQ ID NO:63:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 95 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO-ACID OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 22 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 23..42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 23-42 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 75
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 75 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 76..95
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 76-95 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
   35   40   45

Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   50   55   60

Xaa Xaa Xaa Xaa Xaa Arg Gly Asp Ser Pro Cys Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
          85   90   95

(2) INFORMATION FOR SEQ ID NO:64:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 117 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not-relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACID OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 28
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 28 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 29..48
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 29-48 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 49
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 49 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 70 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 71..90
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 71-90 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 98..117
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 98-117 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Lys Phe Thr Ile Val Phe Cys Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    35   40   45

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
    85   90   95

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Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    100   105   110

Xaa Xaa Xaa Xaa Xaa
    115

(2) INFORMATION FOR SEQ ID NO:65:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 97 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 28
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 28 CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 29..48
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 29-48 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 49
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 49 CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 70 CAN BE TYR OR TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 78..97
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 78-97 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Lys Phe Thr Ile Val Phe Cys Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    35   40   45

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys Val Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    85   90   95

Xaa

(2) INFORMATION FOR SEQ ID NO:66:

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(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 99 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 22 CAN BE TYR, TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 23..42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 23-42 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 51..70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 51-70 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 79
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 79 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 80..99
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa FROM 80-99 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Pro Lys Lys Lys Arg
    35   40   45

Lys Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Glu Ile Leu Asp Val Ser Pro Thr Cys Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    85   90   95

Xaa Xaa Xaa

(2) INFORMATION FOR SEQ ID NO:67:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 78 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa FROM 1-20 CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

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(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 22 CAN BE TYR, TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30..49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 30-49 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 58
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 58 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 59..78
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 59-78 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:67:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Pro Lys Lys Lys Arg Lys Val Xaa Xaa Xaa
      20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  35   40   45

Xaa Glu Ile Leu Asp Val Ser Pro Thr Cys Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  65   70   75

(2) INFORMATION FOR SEQ ID NO:68:

   (i) SEQUENCE CHARACTERISTICS:
      (A) LENGTH: 77 amino acids
      (B) TYPE: amino acid
      (C) STRANDEDNESS: not relevant
      (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 1..20
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACID OR CAN BE
ABSENT"

  (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 21
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

  (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 28
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 28 CAN BE ABSENT"

  (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 29..48
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 29-48 CAN BE ANY AMINO ACID OR CAN BE
ABSENT"

  (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 57
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 57 CAN BE ABSENT"

  (ix) FEATURE:
     (A) NAME/KEY: Modified-site
     (B) LOCATION: 58..77
     (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 58-77 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:68:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Lys Phe Thr Ile Val Phe Cys Xaa Xaa Xaa Xaa
         20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     35   40   45

Glu Ile Leu Asp Val Pro Ser Thr Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75

(2) INFORMATION FOR SEQ ID NO:69:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 93 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 21 CAN BE TYR, TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22..41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 22-41 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 49..68
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 49-68 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 74..93
    (D) OTHER INFORMATION: /product= "OTHER"
/note="Xaa AT POSITIONS 74-93 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:69:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys Val
   35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   50   55   60

Xaa Xaa Xaa Xaa Arg Gly Asp Met Phe Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   85   90

(2) INFORMATION FOR SEQ ID NO:70:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 94 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 1-20 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS AT POSITION 21 CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22..41
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 22-41 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 42 CAN BE ANY AMINO ACID OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 50-69 CAN BE ANY AMINO ACID OR CAN BE
ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 75..94
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITIONS 75-94 CAN BE ANY AMINO ACIDS OR CAN BE
ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:70:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
     20   25   30

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
   35   40   45

Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   50   55   60

Xaa Xaa Xaa Xaa Xaa Arg Gly Asp Met Phe Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    85   90

(2) INFORMATION FOR SEQ ID NO:71:

   (i) SEQUENCE CHARACTERISTICS:
      (A) LENGTH: 94 amino acids
      (B) TYPE: amino acid
      (C) STRANDEDNESS: not relevant
      (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (ix) FEATURE:
      (A) NAME/KEY: Modified-site
      (B) LOCATION: 1..20
      (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

   (ix) FEATURE:
      (A) NAME/KEY: Modified-site
      (B) LOCATION: 21
      (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE TYR, TRP OR CAN BE ABSENT"

   (ix) FEATURE:
      (A) NAME/KEY: Modified-site
      (B) LOCATION: 22..41
      (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

   (ix) FEATURE:
      (A) NAME/KEY: Modified-site
      (B) LOCATION: 49..68
      (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 75..94
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:71:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys Val
   35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
 50   55   60

Xaa Xaa Xaa Xaa Gly Arg Gly Asp Ser Pro Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    85   90

(2) INFORMATION FOR SEQ ID NO:72:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 94 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE TYR, TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 23..42
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 75..94
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:72:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys
   35   40   45

Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa Xaa Arg Gly Asp Ser Pro Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      85   90

(2) INFORMATION FOR SEQ ID NO:73:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 116 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "other"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 27
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 28..47
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 48
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 49..68
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 69
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa AT POSITION 69 CAN BE TYR, TRP, OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 70..89
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 97..116
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:73:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Lys Phe Thr Ile Val Phe Cys Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
    35   40   45

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys Val
    85   90   95

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    100   105   110

Xaa Xaa Xaa Xaa
  115

(2) INFORMATION FOR SEQ ID NO:74:

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 97 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 28
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 29..48
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Active-site
    (B) LOCATION: 49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 70
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE TYR, TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 78..97
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:74:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Lys Phe Thr Ile Val Phe Cys Xaa Xaa Xaa Xaa
     20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   35   40   45

Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Pro Lys Lys Lys Arg Lys Val Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   85   90   95

Xaa

(2) INFORMATION FOR SEQ ID NO:75:

(i) SEQUENCE CHARACTERISTICS:
  (A) LENGTH: 97 amino acids
  (B) TYPE: amino acid
  (C) STRANDEDNESS: not relevant
  (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
  (A) NAME/KEY: Modified-site
  (B) LOCATION: 1..20
  (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(ix) FEATURE:
  (A) NAME/KEY: Modified-site
  (B) LOCATION: 21
  (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE TYR, TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22..41
    (D) OTHER INFORMATION: /product= "other"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 50..69
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 78..97
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:75:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
   20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Pro Lys Lys Lys Arg Lys
   35   40   45

Val Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
  50   55   60

Xaa Xaa Xaa Xaa Xaa Glu Ile Leu Asp Val Ser Pro Thr Xaa Xaa Xaa
65   70   75   80

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    85   90   95

Xaa

(2) INFORMATION FOR SEQ ID NO:76:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 77 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

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(ii) MOLECULE TYPE: peptide

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 1..20
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 21
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 22
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE TYR, TRP OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 30..49
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(ix) FEATURE:
    (A) NAME/KEY: Modified-site
    (B) LOCATION: 58..77
    (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:76:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Cys Xaa Pro Lys Lys Lys Arg Lys Val Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    35   40   45

Xaa Glu Ile Leu Asp Val Ser Pro Thr Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

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Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75

(2) INFORMATION FOR SEQ ID NO:77:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 75 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 27
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "CYS CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 28..47
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID AND CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 56..75
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:77:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Lys Phe Thr Ile Val Phe Cys Xaa Xaa Xaa Xaa Xaa
    20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Glu
   35   40   45

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Ile Leu Asp Val Pro Ser Thr Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70   75

(2) INFORMATION FOR SEQ ID NO:78:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 70 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: circular

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY CATIONIC AMINO ACIDS OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21..40
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 51..70
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACIDS OR CAN BE ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:78:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Glu Ile Leu Asp Val Pro Ser
        35   40   45

Thr Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa
65   70

(2) INFORMATION FOR SEQ ID NO:79:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 71 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: circular

    (ii) MOLECULE TYPE: peptide

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 1..20
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY CATIONIC AMINO ACID OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 21..40
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

    (ix) FEATURE:
        (A) NAME/KEY: Modified-site
        (B) LOCATION: 52..71
        (D) OTHER INFORMATION: /product= "OTHER"
/note= "Xaa CAN BE ANY AMINO ACID OR CAN BE ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:79:

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1   5   10   15

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        20   25   30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys His Ala Ile Arg Gly Asp Thr
        35   40   45

Phe Ala Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    50   55   60

Xaa Xaa Xaa Xaa Xaa Xaa Xaa
65   70

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(2) INFORMATION FOR SEQ ID NO:80:

    (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 14 amino acids
       (B) TYPE: amino acid
       (C) STRANDEDNESS: not relevant
       (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (ix) FEATURE:
       (A) NAME/KEY: Modified-site
       (B) LOCATION: 14
       (D) OTHER INFORMATION: /product= "OTHER"
/note= "G AT POSITION 14 CAN BE ABSENT"

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:80:

Gly Lys Phe Thr Ile Val Phe Asp Asp Asp Asp Asp Asp Gly
1   5   10

(2) INFORMATION FOR SEQ ID NO:81:

    (i) SEQUENCE CHARACTERISTICS:
       (A) LENGTH: 28 amino acids
       (B) TYPE: amino acid
       (C) STRANDEDNESS: not relevant
       (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:81:

Lys Phe Thr Ile Val Phe Gly Gly Gly Leu Phe Glu Ala Ile Ala Glu
1   5   10   15

Phe Ile Glu Gly Gly Trp Glu Gly Leu Ile Glu Gly
          20   25

(2) INFORMATION FOR SEQ ID NO:82:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 27 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:82:

    Gly Leu Phe Glu Ala Ile Ala Glu Phe Ile Glu Gly Gly Trp Glu Gly
    1   5   10   15

    Leu Ile Glu Gly Cys Lys Phe Thr Ile Val Phe
          20   25

(2) INFORMATION FOR SEQ ID NO:83:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 38 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:83:

    Cys Gly Tyr Gly Gly Gly Gly Gly Gly Pro Lys Lys Lys Arg Lys Val
    1   5   10   15

Gly Gly Gly Leu Phe Glu Ala Ile Ala Glu Phe Ile Glu Gly Gly Trp
            20   25   30

Glu Gly Leu Ile Glu Gly
        35

(2) INFORMATION FOR SEQ ID NO:84:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 38 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:84:

    Gly Leu Phe Glu Ala Ile Ala Glu Phe Ile Glu Gly Gly Trp Glu Gly
    1   5   10   15

    Leu Ile Glu Gly Gly Gly Tyr Gly Gly Gly Gly Gly Gly Pro Lys Lys
            20   25   30

    Lys Arg Lys Val Gly Gly
        35

(2) INFORMATION FOR SEQ ID NO:85:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:85:

Lys Phe Thr Thr Ile Val Phe Cys Gly Tyr Gly Pro Lys Lys Lys Arg
1   5   ,   10   15

Lys Val Gly Gly
        20

(2) INFORMATION FOR SEQ ID NO:86:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 19 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:86:

Cys Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly Lys Phe Thr
1   5   10   15

Ile Val Phe

(2) INFORMATION FOR SEQ ID NO:87:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 30 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

(ix) FEATURE:
   (A) NAME/KEY: Modified-site
   (B) LOCATION: 30
   (D) OTHER INFORMATION: /product= "OTHER"
/note= "G AT POSITION 30 CAN BE ABSENT"

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:87:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
   1   5   10   15

Gly Lys Phe Thr Ile Val Phe Asp Asp Asp Asp Asp Asp Gly
      20   25   30

(2) INFORMATION FOR SEQ ID NO:88:

   (i) SEQUENCE CHARACTERISTICS:
      (A) LENGTH: 44 amino acids
      (B) TYPE: amino acid
      (C) STRANDEDNESS: not relevant
      (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

   (iii) HYPOTHETICAL: NO

   (iv) ANTI-SENSE: NO

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:88:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Lys Phe Thr Ile Val Phe Gly Gly Gly Leu Phe Glu Ala Ile Ala Glu
      20   25   30

Phe Ile Glu Gly Gly Trp Glu Gly Leu Ile Glu Gly
   35   40

(2) INFORMATION FOR SEQ ID NO:89:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 43 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:89:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Gly Leu Phe Glu Ala Ile Ala Glu Phe Ile Glu Gly Gly Trp Glu Gly
     20   25   30

Leu Ile Glu Gly Cys Lys Phe Thr Ile Val Phe
     35   40

(2) INFORMATION FOR SEQ ID NO:90:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 54 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:90:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Gly Leu Phe Glu Ala Ile Ala Glu Phe Ile Glu Gly Gly Trp Glu Gly
         20   25   30

Leu Ile Glu Gly Gly Gly Tyr Gly Gly Gly Gly Gly Gly Pro Lys Lys
      35   40   45

Lys Arg Lys Val Gly Gly
   50

(2) INFORMATION FOR SEQ ID NO:91:

(i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 36 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:91:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Lys Phe Thr Thr Ile Val Phe Cys Gly Tyr Gly Pro Lys Lys Lys Arg
         20   25   30

Lys Val Gly Gly
   35

(2) INFORMATION FOR SEQ ID NO:92:

(i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 37 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:92:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Gly Gly Cys Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly Lys
            20   25   30

Phe Thr Ile Val Phe
        35

(2) INFORMATION FOR SEQ ID NO:93:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 56 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:93:

    Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
    1   5   10   15

    Gly Gly Cys Gly Tyr Gly Gly Gly Gly Gly Gly Pro Lys Lys Lys Arg
            20   25   30

    Lys Val Gly Gly Gly Leu Phe Glu Ala Ile Ala Glu Phe Ile Glu Gly
            35   40   45

    Gly Trp Glu Gly Leu Ile Glu Gly
        50   55

(2) INFORMATION FOR SEQ ID NO:94:

186

(i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 24 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:94:

    Ser Ser Asp Asp Glu Ala Thr Ala Asp Ser Gln His Ser Thr Pro Pro
    1   5   10   15

    Lys Lys Lys Arg Lys Val Gly Gly
            20

(2) INFORMATION FOR SEQ ID NO:95:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 40 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:95:

    Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
    1   5   10   15

    Ser Ser Asp Asp Glu Ala Thr Ala Asp Ser Gln His Ser Thr Pro Pro
            20   25   30

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Lys Lys Lys Arg Lys Val Gly Gly
     35   40

(2) INFORMATION FOR SEQ ID NO:96:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 25 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:96:

    Met Ser Tyr Tyr His His His His His His Asp Tyr Asp Ile Pro Thr
    1   5   10   15

    Thr Glu Asn Leu Tyr Phe Gln Gly Ser
          20   25

(2) INFORMATION FOR SEQ ID NO:97:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 37 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:97:

    Met Ser Tyr Tyr His His His His His His Asp Tyr Asp Ile Pro Thr
    1   5   10   15

Thr Glu Asn Leu Tyr Phe Gln Gly Ser Gly Tyr Gly Pro Lys Lys Lys
        20   25   30

Arg Lys Val Gly Gly
   35

(2) INFORMATION FOR SEQ ID NO:98:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 30 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (iii) HYPOTHETICAL: NO

  (iv) ANTI-SENSE: NO

  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:98:

Met Ser Tyr Tyr His His His His His His Asp Tyr Asp Ile Pro Thr
1   5   10   15

Thr Glu Asn Leu Tyr Phe Gln Ser Lys Phe Thr Ile Val Phe
        20   25   30

(2) INFORMATION FOR SEQ ID NO:99:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 44 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (iii) HYPOTHETICAL: NO

  (iv) ANTI-SENSE: NO

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:99:

Met Ser Tyr Tyr His His His His His His Asp Tyr Asp Ile Pro Thr
1   5   10   15

Thr Glu Asn Leu Tyr Phe Gln Ser Gly Leu Phe Glu Ala Ile Ala Glu
        20   25   30

Phe Ile Glu Gly Gly Trp Glu Gly Leu Ile Glu Gly
        35   40

(2) INFORMATION FOR SEQ ID NO:100:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 30 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:100:

Met Ser Tyr Tyr His His His His His His Asp Tyr Asp Ile Pro Thr
1   5   10   15

Thr Glu Asn Leu Tyr Phe Gln Ser Arg Gly Asp Ser Pro Cys
        20   25   30

(2) INFORMATION FOR SEQ ID NO:101:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 10 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:101:

Met Ser Tyr Tyr His His His His His His
1   5   10

(2) INFORMATION FOR SEQ ID NO:102:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 22 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:102:

Met Ser Tyr Tyr His His His His His His Gly Tyr Gly Pro Lys Lys
1   5   10   15

Lys Arg Lys Val Gly Gly
       20

(2) INFORMATION FOR SEQ ID NO:103:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 16 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:103:

Met Ser Tyr Tyr His His His His His His Lys Phe Thr Ile Val Phe
1   5   10   15

(2) INFORMATION FOR SEQ ID NO:104:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 30 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:104:

Met Ser Tyr Tyr His His His His His His Gly Leu Phe Glu Ala Ile
1   5   10   15

Ala Glu Phe Ile Glu Gly Gly Trp Glu Gly Leu Ile Glu Gly
        20   25   30

(2) INFORMATION FOR SEQ ID NO:105:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 16 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:105:

Met Ser Tyr Tyr His His His His His His Arg Gly Asp Ser Pro Cys
1   5   10   15

(2) INFORMATION FOR SEQ ID NO:106:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 6 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (iii) HYPOTHETICAL: NO

  (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:106:

His His His His His His
1   5

(2) INFORMATION FOR SEQ ID NO:107:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 16 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

  (iii) HYPOTHETICAL: NO

  (iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:107:

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Gly Gly Gly Gly Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly
1   5   10   15

(2) INFORMATION FOR SEQ ID NO:108:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 12 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

 (iii) HYPOTHETICAL: NO

 (iv) ANTI-SENSE: NO

 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:108:

Gly Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly
1   5   10

(2) INFORMATION FOR SEQ ID NO:109:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 14 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

 (iii) HYPOTHETICAL: NO

 (iv) ANTI-SENSE: NO

 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:109:

Gly Cys Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly
1   5   10

(2) INFORMATION FOR SEQ ID NO:110:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 9 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:110:

Gly Gly Arg Gly Asp Met Phe Gly Gly
1   5

(2) INFORMATION FOR SEQ ID NO:111:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 22 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:111:

Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly Gly Gly Gly Arg
1   5   10   15

Gly Asp Met Phe Gly Gly
      20

(2) INFORMATION FOR SEQ ID NO:112:

195

(i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:112:

    Gly Leu Phe Gly Ala Ile Ala Gly Phe Ile Glu Asn Gly Trp Glu Gly
    1   5   10   15

    Met Ile Asp Gly
            20

(2) INFORMATION FOR SEQ ID NO:113:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:113:

    Gly Leu Phe Gly Ala Ile Ala Gly Phe Ile Glu Asn Gly Trp Glu Gly
    1   5   10   15

    Leu Ile Glu Gly
            20

196

(2) INFORMATION FOR SEQ ID NO:114:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 20 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:114:

Gly Leu Phe Lys Ala Ile Ala Lys Phe Ile Lys Gly Gly Trp Lys Gly
1   5   10   15

Leu Ile Lys Gly
      20

(2) INFORMATION FOR SEQ ID NO:115:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 12 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS:
        (D) TOPOLOGY: not relevant

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:115:

Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly
1   5   10

(2) INFORMATION FOR SEQ ID NO:116:

197

(i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 31 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:116:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Gly Gly Cys Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly
      20   25   30

(2) INFORMATION FOR SEQ ID NO:117:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 36 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:117:

Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
1   5   10   15

Cys Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly Gly Gly Arg
      20   25   30

198

Gly Asp Ser Pro
            35

(2) INFORMATION FOR SEQ ID NO:118:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 25 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:118:

    Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys Lys
    1   5   10   15

    Gly Gly Arg Gly Asp Ser Pro Cys Gly
            20   25

(2) INFORMATION FOR SEQ ID NO:119:

    (i) SEQUENCE CHARACTERISTICS:
        (A) LENGTH: 25 amino acids
        (B) TYPE: amino acid
        (C) STRANDEDNESS: not relevant
        (D) TOPOLOGY: linear

    (ii) MOLECULE TYPE: peptide

    (iii) HYPOTHETICAL: NO

    (iv) ANTI-SENSE: NO

    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:119:

    Gly Arg Gly Asp Ser Pro Cys Gly Gly Lys Lys Lys Lys Lys Lys Lys
    1   5   10   15

Lys Lys Lys Lys Lys Lys Lys Lys Lys
        20   25

(2) INFORMATION FOR SEQ ID NO:120:

  (i) SEQUENCE CHARACTERISTICS:
    (A) LENGTH: 22 amino acids
    (B) TYPE: amino acid
    (C) STRANDEDNESS: not relevant
    (D) TOPOLOGY: linear

  (ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:120:

Cys Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly Gly Gly Arg
1   5   10   15

Gly Asp Ser Pro Cys Gly
        20

200

Lys Lys Lys Lys L's Lys Lys Lys Lys
      20   25

(2) INFORMATION FOR SEQ ID NO:120:

   (i) SEQUENCE CHARACTERISTICS:
     (A) LENGTH: 22 amino acids
     (B) TYPE: amino acid
     (C) STRANDEDNESS: not relevant
     (D) TOPOLOGY: linear

   (ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

   (xi) SEQUENCE DESCRIPTION: SEQ ID NO:120:

Cys Gly Tyr Gly Pro Lys Lys Lys Arg Lys Val Gly Gly Gly Gly Arg
1   5   10   15

Gly Asp Ser Pro Cys Gly
         20